## Amendments to the Claims

of said hybrid X1069G having been deposited under ATCC Accession number
Claim 2 (Currently amended): A maize plant, or its parts a part thereof, produced by growing the seed of claim 1.
Claim 3 (Original): Pollen of the plant of claim 2.
Claim 4 (Original): An ovule of the plant of claim 2
Claims 5-62 (Canceled)
Claim 63 (Previously presented): A tissue culture of regenerable cells produced from the plant of claim 2.
Claim 64 (Previously presented): Protoplasts produced from the tissue culture of claim 63.
Claim 65 (Currently amended): The tissue culture produced from the plant of claim 263, wherein cells of the tissue culture are from a tissue selected from the group consisting of leaf, pollen, embryo, root, root tip, anther, silk, flower, kernel, ear, cob, husk and stalk.
Claim 66 (Previously presented):  A maize plant regenerated from the tissue culture of claim 63, said plant having all the morphological and physiological characteristics of hybrid maize plant X1069G, representative seed of said plant having been deposited under ATCC Accession No
Claim 67 (Previously presented): A method for producing an FI hybrid maize seed, comprising crossing the plant of claim 2 with a different maize plant and harvesting the resultant F1 hybrid maize seed.

Claim 68 (Previously presented):	A method of prod	lucing a male sterile hy	brid maize plant
comprising transforming at least one	of inbred maize p	arent plants GE535769	and GE515721,
representative samples of which have	been deposited a	s and	_respectively, with
a nucleic acid molecule that confers n	nale sterility and	prossing said inbred ma	aize parent plants to
produce said male sterile hybrid maiz	e plant.		
Claim 69 (Previously presented):	A male sterile ma	ize hybrid plant produ	ced by the method
of claim 68.			
Claim 70 (Previously presented):	A method of prod	lucing an herbicide res	istant hybrid maize
plant comprising transforming at least	t one of inbred m	aize parent plants GE5:	35769 and
GE515721, representative samples of	which have been	deposited as	and
respectively, with a transgene that cor	nfers herbicide re	sistance to generate an	herbicide resistant
inbred maize parent plant and crossing	g said inbred mai	ze parent plants to proc	luce said herbicide
resistant hybrid maize plant,		•	
Claim 71 (Previously presented):	An herbicide resi	stant hybrid maize plan	it produced by the
method of claim 70.			•
Claim 72 (Previously presented):	The herbicide res	 istant hybrid maize pla	nt of claim 71,
wherein the transgene confers resistan	ce to an herbicid	selected from the gro	up consisting of:
imidazolinone, sulfonylurea, glyphosa	ate, glufosinate, L	-phosphinothricin, trla:	zine and
benzonitrile.			
Claim 73 (Previously presented):	A method of prod	lucing an insect resistar	nt hybrid maize
plant comprising transforming at least	one of inbred ma	aize parent plants GE53	35769 and
GE515721, representative samples of	which have been	deposited as	and
respectively, with a transgene that con			
maize parent plant and crossing said in		_	
hybrid maize plant.	•		

Claim 74 (Previously presented): claim 73.	An insect resistar	nt maize plant produced by the method of
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Claim 75 (Currently amended):	The insect resists	nt maize plant of claim 74, wherein the
transgene comprises a transgene enco	ding encodes a B	acillus thuringiensis endotoxin.
Claim 76 (Previously presented):	A method of proc	lucing a disease resistant hybrid maize
plant comprising transforming at least	-	•
GE515721, representative samples of		
, .	•	tance to generate a disease resistant inbred
		nt plants to produce said disease resistant
hybrid maize plant.	nored marze pare	in plants to produce and disease resistant
nyone maize plant.		
Claim 77 (Previously presented):	A disease resistar	it hybrid maize plant produced by the
method of claim 76.	11 discuso resistar	is riyond maze plant produced by the
mediod of claim 70.		
Claim 78 (Previously presented):	A method of proc	lucing a hybrid maize plant with decreased
phytate content comprising transform	ing at least one of	inbred maize parent plants GE535769
and GE515721, representative sample	es of which have I	peen deposited as and
		rate an inbred maize parent plant with
decreased phytate content and crossin	g said inbred mai	ze parent plants to produce said hybrid
maize plant that confers decreased phy	ytate content.	
	-	
Claim 79 (Previously presented):	A hybrid maize p	ant with decreased phytate content
produced by the method of claim 78.		
Claim 80 (Previously presented):	A method of proc	ucing a hybrid maize plant with modified
fatty acid metabolism or modified car	bohydrate metabo	lism comprising transforming at least one
of inbred maize parent plants GE5357	769 and GE51572	1, representative samples of which have
been deposited as and	respectively;	with a transgene encoding a protein
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selected from the group consisting of stearyl-ACP desaturase, fructosyltransferase, levansucrase, alpha-amylase, invertase and starch branching enzyme to generate an inbred maize parent plant with modified fatty acid metabolism or modified carbohydrate metabolism and crossing said inbred maize parent plants to produce said hybrid maize plant that confers modified fatty acid metabolism or modified carbohydrate metabolism.

inbred maize parent plants to produce said hybrid maize plant that confers modified fatty acid
metabolism or modified carbohydrate metabolism.
Claim 81 (Previously presented): A hybrid maize plant produced by the method of claim 80.
Claim 82 (Previously presented): The hybrid maize plant of claim 81 wherein the transgene
confers a trait selected from the group consisting of waxy starch and increased amylose starch.
Claim 83 (Previously presented): A maize plant, or part thereof, having all the physiological
and morphological characteristics of the hybrid maize plant X1069G, representative seed of said
plant having been deposited under ATCC Accession No
Claim 84 (Currently amended): A method of introducing a desired trait into a hybrid maize
line X1069G comprising:
(a) crossing at least one of inbred maize parent plants GE535769 and GE515721,
representative samples of which have been deposited under ATCC Accession Nos. as
and respectively, with another maize line that comprises a desired trait, to produce F1
progeny plants, wherein the desired trait is selected from the group consisting of male sterility,
herbicide resistance, insect resistance, disease resistance and waxy starch;
(b) selecting said F1 progeny plants that have the desired trait to produce selected F1
progeny plants;
(c) backcrossing the selected progeny plants with said inbred maize parent plant to
produce backcross progeny plants;
(d) selecting for backcross progeny plants that have the desired trait and morphological
and physiological characteristics of said inbred maize parent plant;

times in succession to produce selected fourth or higher backcross progeny plants;

(e) repeating the steps of backcrossing to said inbred maize parent plant three or more

(f) crossing said backcross progeny plant with the other inbred maize parent plant to generate a hybrid maize line X1069G with the desited trait and all of the morphological and physiological characteristics of hybrid maize line X1069G listed in Table 1 as determined at-athe 5% significance level when grown in the same environmental conditions.

Claim 85 (Currently amended): A plant produced by the method of claim 84, wherein the plant has the desired trait and all of the physiological and morphological characteristics of hybrid maize line X1069G listed in Table 1 as determined at the 5% significance level when grown in the same environmental conditions.

Claim 86 (Previously presented): The plant of claim 85 wherein the desired trait is herbicide resistance and the resistance is conferred to an herbicide selected from the group consisting of: imidazolinone, sulfonylurea, glyphosate, glufosinate, L-phosphinothricin, triazine and benzonitrile.

Claim 87 (Previously presented): The plant of claim 85 wherein the desired trait is insect resistance and the insect resistance is conferred by a transgene encoding a *Bacillus thuringiensis* endotoxin

Claim 88 (Previously presented): The plant of claim 85 wherein the desired trait is male sterility and the trait is conferred by a cytoplasmic nucleic acid molecule that confers male sterility.

Claim 89 (Currently amended): A method of <u>modifying introducing modified</u> fatty acid metabolism, <del>modified</del> phytic acid metabolism or <del>modified</del> carbohydrate metabolism <del>into</del>in a hybrid maize line X1069G comprising:

(a) crossing at least one of inbred maize parent plants GE535769 and GE515721, representative samples of which have been deposited <u>under ATCC Accession Nos.</u> as and \_\_\_\_\_\_ respectively, with another maize line that comprises a desired trait nucleic acid molecule encoding an enzyme-to-produce F1 progeny plants, wherein the desired trait is selected

from the group consisting of phytase, stearyl-ACP desaturase, fructosyltransferase, levansucrase, alpha-amylase, invertase and starch branching enzyme

- (b) selecting said F1 progeny plants that have the desired trait said nucleic acid molecule to produce selected F1 progeny plants;
- (c) backcrossing the selected progeny plants with said inbred maize parent plant to produce backcross progeny plants;
- (d) selecting for backcross progeny plants that have the desired trait said nucleic acid molecule and morphological and physiological characteristics of said inbred maize parent plant;
- (e) repeating the steps of backcrossing to said inbred maize parent plant three or more times in succession to produce selected fourth or higher backcross progeny plants;
- (f) crossing said backcross progeny plant with the other inbred maize parent plant to generate a hybrid maize line X1069G with the desired trait that comprises said nucleic acid molecule and has all of the morphological and physiological characteristics of hybrid maize line X1069G listed in Table 1 as determined at athe 5% significance level when grown in the same environmental conditions.

Claim 90 (Currently amended): A plant produced by the method of claim 89, wherein the plant has modified fatty acid metabolism, modified phytic acid metabolism or modified carbohydrate metabolism and all of the physiological and morphological characteristics of hybrid maize line X1069G listed in Table 1 as determined at athe 5% significance level when grown in the same environmental conditions.

Claim 91 (New): A method for producing a maize seed, comprising crossing the plant of claim 2 with itself or a different maize plant and harvesting the resultant maize seed.